

Table S2. *Number of reads for each sample analyzed by RNA-sequencing*

Sample	Medium	Reads
Met_neg_1	BMM	803313
Met_neg_2	BMM	726389
Met_neg_3	BMM	596414
Met_pos_1	BMM + Methionine	696835
Met_pos_2	BMM + Methionine	536542
Met_pos_3	BMM + Methionine	756652